## SEQUENCE LISTING

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<!!?> Clemiennen, Stephanie K.
      Schuster, Debra K.
· 124 · CIRI HOMOLOGUE FROM MELON
.1*0 - 4357-0929.30
- 140 - Not Yet Assigned
-141 - Filed Herewith
- 150 ⋅ H3 60/218,307
×151 × 2000-07-14
160 > 8
+170 > FastSEQ for Windows Version 4.0
3.200 × 1
+211+ 3236
\mathrm{AUCL} + \mathrm{DMA}
<213 · Capumis melo</pre>
× 520 ×
.lil . musc_feature
(31)...(81
-203 + r = A,T,C or G
91100 ×
+2.1 + misc_feature
- 592 · (593) · · · 5 · 4)
+203+r. = A,T,C \rightarrow c G
\sim 400 \times 1
 uttogattgt ogggangaga gagoagaaaa ttaaaaccag aatotocaac acacaaacct
                                                                           60
 tocaccectt caacaatgge nggattetag ggtttetatg ggtttaaggt gatacagttt
                                                                           120
 octaatitet coatgaaat geetggaegg aggtetgatt actetettt aagteaaatt
                                                                           150
 reggacyang aggreggacy gragotted acttentitt acquetcogt agragetggg
                                                                           240
 ggaaacgtta tcaaagggag aaccgatagg gtttttgatt gggatgggag tggtgatcac
                                                                           300
 aggttalada ogcaggogta toggataggg aacctgtatt catggattgg titacagaga
                                                                           360
 cattocaying gaageagota egatgatago teteteteta gigattaeta egeacegaeg
                                                                           4_0
 rtatrauseo etgesjesas tgagateast gesttggast stateetegs tgatgattte
                                                                           430
 equatigatga aagetgtggg aagtggaggt tegtetggaa agagetggge esageagaeg
                                                                           5.40
 guagagaget troagttgea geagedettg gttettagge titetteaga tgmmaettgt
                                                                           F. 00
 geogatgate coaactttat ggatocgatt coagacgagg cagetttaag atogitatig
                                                                           4.60
 attroagety aggoratote gratogotto toggtaaato gatgcatoto atatroggag
                                                                           720
 assatty: Day stygettetta totaattoat gggatggaco catatgtatg gtoatratgo
                                                                           780
 admaat stgs aagadgatgg gogtatacca toatttgaat ototgaaaac agttgattoc
                                                                           840
 agnatoggti satcaattga agtagttttg atagategge atagtgatge tagstsaaaa
                                                                           90)
 quactquaaa auagagtgca taatatttot tocagttgtg taaccacaaa agaggttgua
                                                                           960
 jatcatatag :@@aqctggt atgcaatcac ttggggggtt cagtttctga gggadaagat
                                                                          1020
 gasttgyttt stysstggaa ggaatgcago gatgasttaa aggaatgttt gggaturgst
                                                                          1030
 qtgattocs: tatgcagott atotgttggc otttgtagac atogtgctct tttarr saaa
                                                                          1140
 gtoptagot; antomatiga tittaccotgt ognatigoda maggatigina atatriplant
                                                                          1200
 agagatgat i inhostottg cottgttagg ttogggettg atagggaata totosi gat
                                                                          1260
  stgattggga ggreaggttg sttatgccaa cotgattstt tgotcaatgg tosat af so
                                                                          1520
 atotoaatti mitoaccatt gogatttosa agadtaaaad otattgaato tabmanigat
                                                                          1380
  ttdaggtbar i rgccaaada gtatttettg gatagecaat caettaatet tgtarii gat
                                                                          1440
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1500

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ccattaaata ggaajga ji agatqgaaaa accatagtgg reactggtga caaggaCaga
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aattotoagt tattaaaria aaaaqoagoo caactgaara ottaagatgg aaagtotgag
                                                                       1620
caatttagat datgrightig i tectocatat agtgtadagi ligaccoctti tgtagaadus
                                                                        1880
gtagtopott täägimatat otoacabatt ggttotgalig attriggagba totottagda
                                                                       1710
tiginicate caaqqatqqa teatqitaas aattiaccat tigiteatqq tagicaqii;
                                                                       1 - 00
attaqaaaac caaargaget ttooottggo ttaqaaqatt tqgttattoo atggacaga:
                                                                       1:50
cttgatttga gggagaaaat tggagcaggt tottttggga otgfatatog tggtgagtgg
                                                                       1920
catggatety atgittgetyt gaagitests acagaacaag acticcatos tgaacgigt.
                                                                       1-1-5()
                                                                       > 40
aatgasttto tgagagaggf tgotatcatg aaatctttac gabatcotas tattgtactg
tttatgggtg oggtgaccaa godaccaaab ttgtccattg tcaccgaati tctatcgaga
                                                                       2.00
                                                                       2.60
ggtayottgt ataggetttt geataagtea ggtgteaaag acatagatga aacaegteja
ataaatangg ottitigatgi ggcaaaggga atgaactaco toracajacg igatooto:a
                                                                       2.120
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                                                                       2.440
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gotgjaacad otqaatggat ggbabbagaa qtabbabgcg abgaacbatb aaatgaaaag
                                                                       -1.7(10)
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tytaatotaa abboagotoa gyttytogoa qotyttyggat ttaaggycaa aaggottyuo
atoccacging angua and casalinger toottaatag togethering goodgangag
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cratggaaas gtoottstit ticcagcatt atggaaacet igaaaccaat gactaaacaa
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gegecacets assassyted cacagacaes etetegetta tytgacaaty tytgtatest
                                                                        2.160
aggaatgeet gabgettigg agggetaagt ubggtacest thaagagatg tetggeatgt
ttaaaaaccat actocaaaca agcaagcacc tgtgctcgta godaaattit ccattgctag
                                                                        - )
tagttacaat tttpaageta agttoettgt accgtgette traagttttt gtgaacggat
                                                                        - - - 1
ggggaagtgt tggaaaastt caactgotag asgattscat caaatttatt stcagttsat
                                                                        \geq \pm 10
aatcatcaaa atgtagagag attataaaaa tgtggatcac ttcatagtcc acaatcaagg
aagtttotad ottitigota tygtgatgaa gaaadttoaa otobatgtoa oodtatttoa
                                                                        3960
                                                                        3120
etteapapat tatttgtttg tatatetatt jgtottacet ttgaggaceg gaseaggaae
taatttigta tatactagtg atcagttgtg gatggatgca atcatgtctt cagtcagast
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 tggtgtttgc tagggaaata tmattgttgt tatttaacag coacttcaaa cattcaatta
                                                                        :240
                                                                        2286
 atitteaccg agretattat tetaaaaaaa aaaaaaaa aaaaaa
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4.2100 £
+ 2110 350
+212: FRT
+213: Cucumis melo
H2200
<!D21> VARIANT
< 222 > (154) \dots (154)
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<!!!! Ama = Any Amano Acid</pre>

<400 > 2

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                                      10
1
Pro Asp Glu Glu Val Gly Thr Gly Ala Ser Thr Ser Phe Tyr Asp Ser
                                  25
             2()
Val Ala Ala Gly Gly Asn Val Ile Lys Gly Arg Thr Asp Arg Val Phe
                                                   45
                             40
Asp Trp Asp Gly Ser Gly Asp His Arg Leu Asn Thr Gln Ala Tyr Arg
                                              \tilde{\mathbf{r}}(t)
                         5.5
    4,11
Ile Gly Asn Leu Tyr Ser Trp Ile Gly Leu Gln Arg His Ser Ser Gly
                                          75
                     70
65
Ser Ser Tyr Asp Asp Ser Ser Leu Ser Ser Asp Tyr Tyr Ala Pro Thr
                                                           35
                                      90
                 85
Leu Ser Asn Pro Ala Ala Asn Glu Ile Asn Ala Leu Glu Tyr Ile Leu
                                                       110
                                  105
             100
Asp Asp Asp Phe Arg Val Met Lys Ala Val Gly Ser Gly Gly Ser Ser
                                                   125
                              120
         115
Gly Lys Ser Trp Ala Gln Gln Thr Glu Glu Ser Phe Gln Leu Gln Gln
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                          135
     130
Pro Leu Val Leu Arg Leu Ser Ser Asp Mad Thr Cys Ala Asp Asp Pro
                                           155
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150

Asn Phe Met Asp Ero lle Pro Asp Glu Ala Ala Den Arg Ser Leu Ser The Ser Ala Glu Ala The Ser His Arg Phy Tri Val Ash Gly Cys Met 190 185 180 Ser Tyr Leu Glu Lys Val Pro Asp Gly Phe Tyr Leu Ile His Gly Met 200 Asp Pro Tyr Val Trp Ser Len Cys Thr Ash Leu Gln Glu Asp Gly Arg 195 215 Ile Pro Ser Phe Glu Ser Leu Lys Thr Val Asp Ser Ser Ile Gly Ser 2:30 Ser Ile Glu Val Val Leu Ile Asp Arg His Ser Asp Ala Ser Leu Lys 250 255 2:45 Glu Leu Gln Asn Arg Val His Asn Ile Ser Ser Ser Cys Val Thr Thr 265 270 260 Lys Glu Val Ala Asp His Ile Ala Lys Leu Val Cys Asn His Leu Gly 275 285 Gly Ser Val Ser Glu Gly Glu Asp Asp Leu Val Ser Ala Trp Lys Glu 3(4) 295 Cys Ser Asp Asp Neu Lys Glu Cys Leu Gly Ser Ail Val Ile Pro Leu 315 310 Cys Ser Leu Ser Val Gly Leu Cys Arg His Arg Ata Leu Leu Phe Lys 33 ( 335 325 Val Leu Ala Asp Ser Ile Asp Leu Pro Cys Arg I.e Ala Lys Gly Cys 350 340 Lys Tyr Cys Thr Arg Asp Asp Ala Ser Ser Cys Leu Val Arg Phe Gly 355 360 365 Leu Asp Arg Glu Tyr Leu Ile Asp Leu Ile Gly Arg Fro Gly Cys Leu 3,80 375 Cys Gln Pro Asp Ser Leu Leu Asn Gly Pro Ser Ser Ile Ser Ile Ser 395 390 Ser Pro Leu Arg Phe Pro Arg Leu Lys Pro Ile Clu Ser Thr Ile Asp 405 Fhe Arg Ser Leu Ala Lys Gln Tyr Phe Leu Asp Cer Gln Ser Leu Asn 425 430 420 Leu Val Phe Asp Glu Ala Ser Ser Gly Asc Val Val Ser Gly Lys Asp 445440 Ala Ala Phe Ser Val Tyr Gln Arg Pro Leu Asn Arg Lys Asp Val Asp 450 455 480 Gly Lys Thr Ile Val Val Thr Gly Asp Lys Asp Arg Asn Ser Gln Leu 475 470 Leu Asn Lys Lys Ala Ala Gln Leu Asn Thr Gln Asp Gly Lys Ser Glu 485 Gln Phe Arg Ser Cys Val Ala Ser Fro Tyr Ser Val Gln Ser Thr Pro 510 500 505 Phe Val Glu Asn Val Val Pro Leu Ser H.s Ilo Ser His Ile Gly Ser 525 520 Glu Asp Ser Glo His Leu Leu Ala Leu Ser His Pro Arg Met Asp His 530 535 Val Asn Asn Leu Pro Phe Val His Gly Ser Gin Leu Ile Arg Lys Pro 550 Asn Glu Leu Ser Leu Gly Leu Glu Asp Leo Val Ile Pro Trp Thr Asp 5.70 565 Leu Asp Leu Arg Glu Lys Ile Gly Ala Gly Jer Phe Gly Thr Val Tyr 535 Arg Gly Glu Trp His Gly Ser Asp Val Ala Cal Lys Ile Leu Thr Glu 605 595 600 Gin Asp Phe His Pro Glu Arg Val Asn Clos En- Leu Arg Glu Val Ala 620 610 615 The Met Lys Ser Leu Arg His Pro Ash Il- 1: Leu Phe Met Gly Ala 630 Val Thr Lys Pro Pro Asn Leu Ser Ile Val The Glu Tyr Leu Ser Arg

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Gly	3er	Leu	Tyr 660	At 1	1,000	Leu	His	Lys 665	Ser	0.9	17:1	Lys	Asp 670	Ile	Asp		
			Arg						Asp								
							Pro		Val								
						Lys			Thr								
					Lys				Phe 730								
									Glu								
				Ser				Ser	Phe								
			Leu				Trp	Cys	Asn								
						Lys	Gly								Asp 800		
					Ala	Ser											
				y Pro	ser				: Ile	Met					Pro		
Мед	Thr	Lys	820 s Glr	n Ala	a Erc	) Pro	Glr 840	ı Glr	n Ser	Arg	Thr	Asp 845	Thr	Leu	ı Ser		
Val	. Met 85(		)				0.10	,									
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	450	5.			~+ ~=	. ++0	C										24
2	tytt	tagg	ga to	ugat	y Tya	ttc	_										
	210> 211:																

<212. DNA <213> Artificial Gequence	
<pre> &lt;22.0&gt; &lt;.200 synthetic oligonarleotide </pre>	
k.;)no. 6 tit.ggtggct tggtcaccgc accc	24
+ 11 0+ 7 + 110+ 24 + 110+ DNA + 113+ Artificial Sequence	
- 120 - - 123 - synthetic oligonucleotide	
0.400 + 7 cutgrightigo tictocatat agitg	24
-210 · 8 -311 · 24 -312 · DNA -313 · Artificial Sequence	
0.200 · 0.203 · synthetic oligonucleotide	
· 40() · 8 teatstatgt etttgacaes tgas	24

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